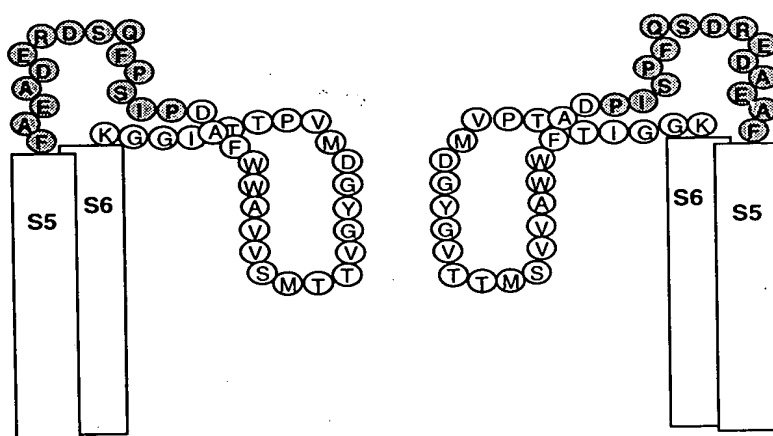


FIG. 1

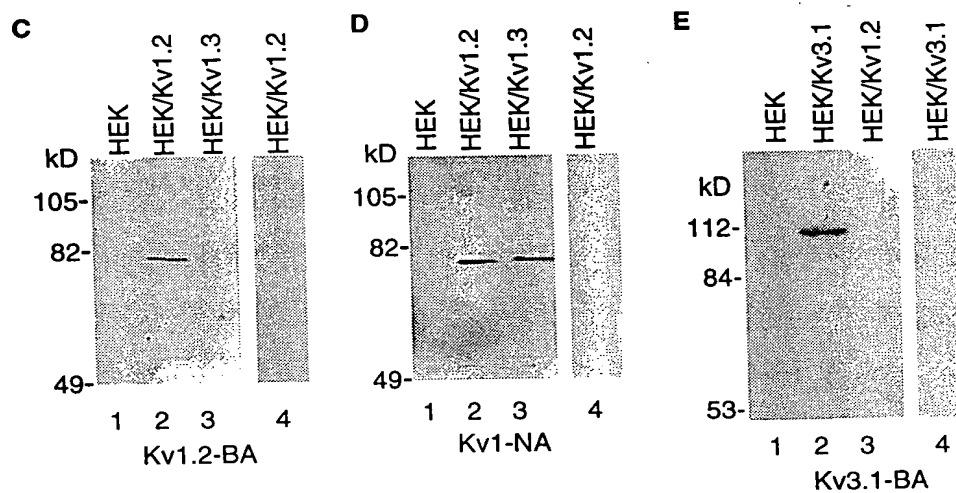
A

		FAE	ADERDSQFPSIP	P region	PTTIGGK	
Kv1.2	S5			DAFWWAVVSM TTVGYGDMV		S6
Kv1.3		---	--DPT-G-S---	-----T-----H	-V-----	
Kv3.1		Y--	<u>RIGAQPNDPSASEH</u> TH-KN--	G-----T-----L-----	-Q-WS-M	

B



**FIG.1**



**FIG. 2**

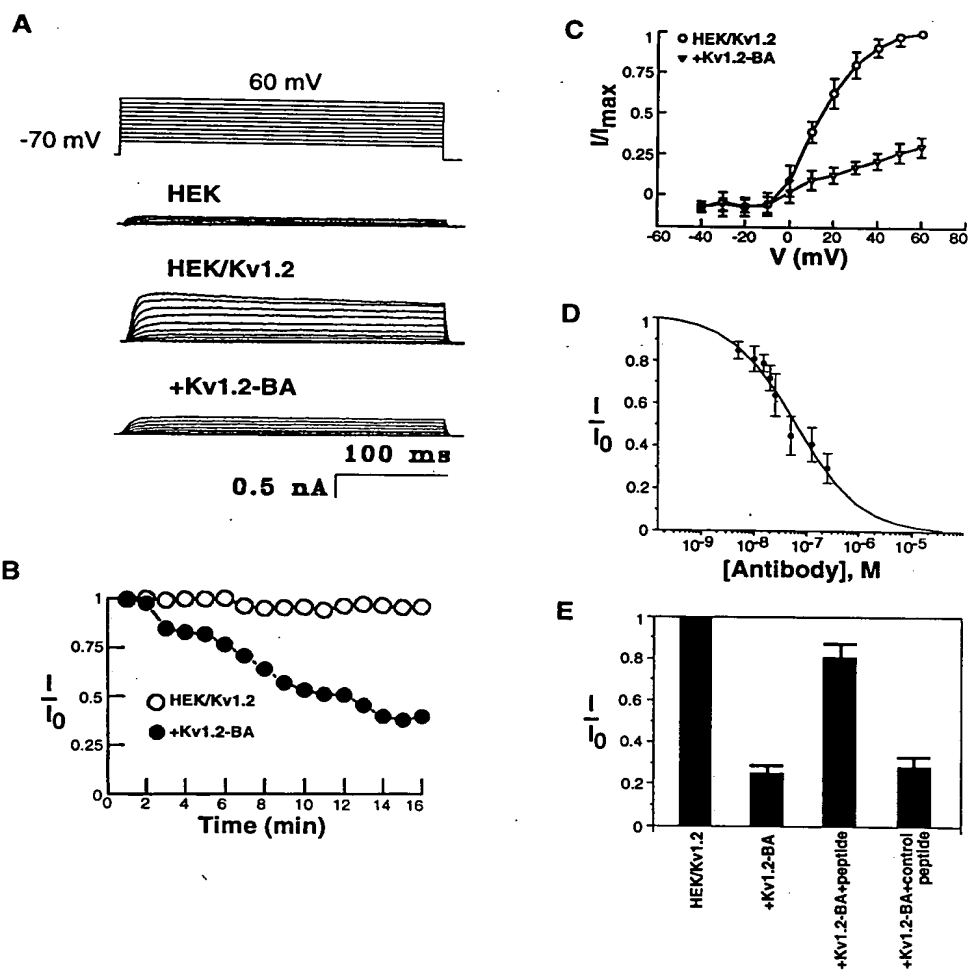


FIG. 3

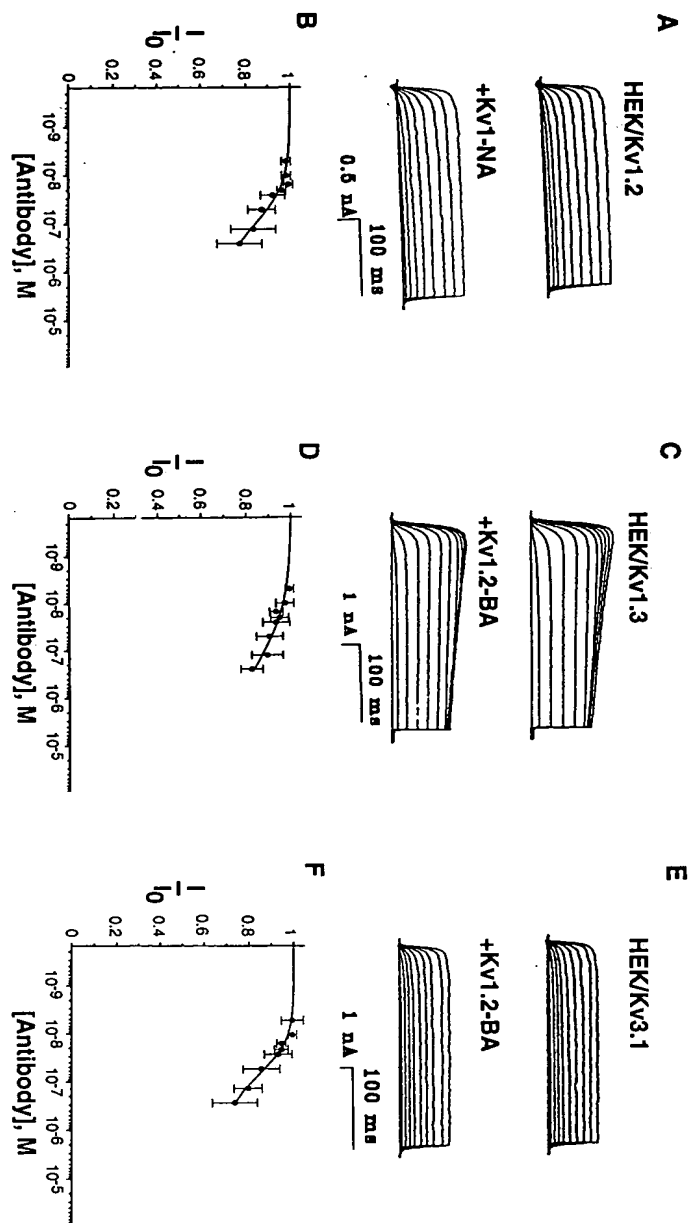
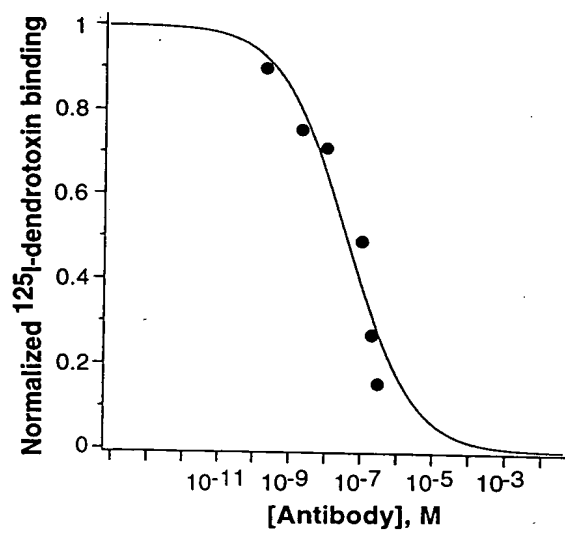


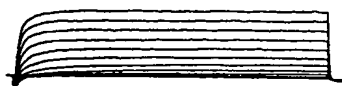
FIG. 4



**FIG. 5**

**A**

**NG108-15**

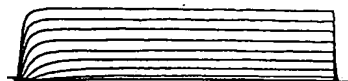


**+Kv1.2-BA**



**B**

**NG108-15**



**+Kv1-NA**



**1 nA** **100 ms**

FIG. 6

